

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/585,923
Source: 1 Fw0
Date Processed by STIC: 2/2/07

ENTERED



IFWO

RAW SEQUENCE LISTING

DATE: 02/02/2007

PATENT APPLICATION: US/10/585,923

TIME: 09:00:52

Input Set : A:\01-18-07 seq list_ST25.txt

Output Set: N:\CRF4\02022007\J585923.raw

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3 <110> APPLICANT: Brion, Aurelie
4      Leissner, Philippe
5      Berard, Cecile
7 <120> TITLE OF INVENTION: Thrombosis Diagnosis/Prognosis Method
9 <130> FILE REFERENCE: 128639
11 <140> CURRENT APPLICATION NUMBER: US 10/585,923
12 <141> CURRENT FILING DATE: 2006-07-13
14 <150> PRIOR APPLICATION NUMBER: PCT/FR05/50083
15 <151> PRIOR FILING DATE: 2005-02-10
17 <160> NUMBER OF SEQ ID NOS: 25
19 <170> SOFTWARE: PatentIn version 3.4
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 23
23 <212> TYPE: DNA
24 <213> ORGANISM: Homo sapiens
26 <400> SEQUENCE: 1
27 aaattctcag aatttctgaa agg                                23
30 <210> SEQ ID NO: 2
31 <211> LENGTH: 22
32 <212> TYPE: DNA
33 <213> ORGANISM: Homo sapiens
35 <400> SEQUENCE: 2
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39 <210> SEQ ID NO: 3
40 <211> LENGTH: 22
41 <212> TYPE: DNA
42 <213> ORGANISM: Homo sapiens
44 <400> SEQUENCE: 3
45 agtgcttaac aagaccatac ta                                    22
48 <210> SEQ ID NO: 4
49 <211> LENGTH: 24
50 <212> TYPE: DNA
51 <213> ORGANISM: Homo sapiens
53 <400> SEQUENCE: 4
54 aacagacctg gaatttgaaa ctaa                                24
57 <210> SEQ ID NO: 5
58 <211> LENGTH: 20
59 <212> TYPE: DNA
60 <213> ORGANISM: Homo sapiens
62 <400> SEQUENCE: 5
63 attactggct cttctgagc                                       20
66 <210> SEQ ID NO: 6
67 <211> LENGTH: 20

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68 <212> TYPE: DNA
69 <213> ORGANISM: Homo sapiens
71 <400> SEQUENCE: 6
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75 <210> SEQ ID NO: 7
76 <211> LENGTH: 20
77 <212> TYPE: DNA
78 <213> ORGANISM: Homo sapiens
80 <400> SEQUENCE: 7
81 tctagaaaca gttgcctggc                20
84 <210> SEQ ID NO: 8
85 <211> LENGTH: 20
86 <212> TYPE: DNA
87 <213> ORGANISM: Homo sapiens
89 <400> SEQUENCE: 8
90 ctaccagcgt gccaccaggt                20
93 <210> SEQ ID NO: 9
94 <211> LENGTH: 16
95 <212> TYPE: DNA
96 <213> ORGANISM: Homo sapiens
99 <220> FEATURE:
100 <221> NAME/KEY: misc_feature
101 <222> LOCATION: (13)..(13)
102 <223> OTHER INFORMATION: n is inosine
104 <400> SEQUENCE: 9
W--> 105 ctggacaggc gangaa                16
108 <210> SEQ ID NO: 10
109 <211> LENGTH: 16
110 <212> TYPE: DNA
111 <213> ORGANISM: Homo sapiens
114 <220> FEATURE:
115 <221> NAME/KEY: misc_feature
116 <222> LOCATION: (13)..(13)
117 <223> OTHER INFORMATION: n is inosine
119 <400> SEQUENCE: 10
W--> 120 ctggacaggc aangaa                16
123 <210> SEQ ID NO: 11
124 <211> LENGTH: 18
125 <212> TYPE: DNA
126 <213> ORGANISM: Homo sapiens
128 <400> SEQUENCE: 11
129 actctcagca agcctcaa                18
132 <210> SEQ ID NO: 12
133 <211> LENGTH: 18
134 <212> TYPE: DNA
135 <213> ORGANISM: Homo sapiens
138 <220> FEATURE:
139 <221> NAME/KEY: misc_feature
140 <222> LOCATION: (13)..(13)

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141 <223> OTHER INFORMATION: n is inosine
143 <400> SEQUENCE: 12
W--> 144 actctcagcg agnctcaa 18
147 <210> SEQ ID NO: 13
148 <211> LENGTH: 51
149 <212> TYPE: DNA
150 <213> ORGANISM: Artificial Sequence
152 <220> FEATURE:
153 <223> OTHER INFORMATION: T7 Phage Polymerase Promoter
155 <400> SEQUENCE: 13
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159 <210> SEQ ID NO: 14
160 <211> LENGTH: 51
161 <212> TYPE: DNA
162 <213> ORGANISM: Artificial Sequence
164 <220> FEATURE:
165 <223> OTHER INFORMATION: T7 Phage Polymerase Promoter
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171 <210> SEQ ID NO: 15
172 <211> LENGTH: 20
173 <212> TYPE: DNA
174 <213> ORGANISM: Homo sapiens
176 <400> SEQUENCE: 15
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180 <210> SEQ ID NO: 16
181 <211> LENGTH: 21
182 <212> TYPE: DNA
183 <213> ORGANISM: Homo sapiens
185 <400> SEQUENCE: 16
186 ctggaaccaa tcccgtgaaa g 21
189 <210> SEQ ID NO: 17
190 <211> LENGTH: 18
191 <212> TYPE: DNA
192 <213> ORGANISM: Homo sapiens
194 <400> SEQUENCE: 17
195 actctcagca agcctcaa 18
198 <210> SEQ ID NO: 18
199 <211> LENGTH: 15
200 <212> TYPE: DNA
201 <213> ORGANISM: Homo sapiens
203 <400> SEQUENCE: 18
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207 <210> SEQ ID NO: 19
208 <211> LENGTH: 44
209 <212> TYPE: DNA
210 <213> ORGANISM: Artificial Sequence
212 <220> FEATURE:
213 <223> OTHER INFORMATION: T7 Phage Polymerase Promoter

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215 <400> SEQUENCE: 19
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219 <210> SEQ ID NO: 20
220 <211> LENGTH: 28
221 <212> TYPE: DNA
222 <213> ORGANISM: Homo sapiens
225 <220> FEATURE:
226 <221> NAME/KEY: misc_feature
227 <222> LOCATION: (19)..(19)
228 <223> OTHER INFORMATION: n is inosine
230 <400> SEQUENCE: 20
W--> 231 cgatcgctgg acaggcgang aacgatcg      28
234 <210> SEQ ID NO: 21
235 <211> LENGTH: 28
236 <212> TYPE: DNA
237 <213> ORGANISM: Homo sapiens
240 <220> FEATURE:
241 <221> NAME/KEY: misc_feature
242 <222> LOCATION: (19)..(19)
243 <223> OTHER INFORMATION: n is inosine
245 <400> SEQUENCE: 21
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249 <210> SEQ ID NO: 22
250 <211> LENGTH: 30
251 <212> TYPE: DNA
252 <213> ORGANISM: Homo sapiens
254 <400> SEQUENCE: 22
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258 <210> SEQ ID NO: 23
259 <211> LENGTH: 30
260 <212> TYPE: DNA
261 <213> ORGANISM: Homo sapiens
264 <220> FEATURE:
265 <221> NAME/KEY: misc_feature
266 <222> LOCATION: (19)..(19)
267 <223> OTHER INFORMATION: n is inosine
269 <400> SEQUENCE: 23
W--> 270 cggtcgactc tcagcgagnc tcaacgaccg      30
273 <210> SEQ ID NO: 24
274 <211> LENGTH: 30
275 <212> TYPE: DNA
276 <213> ORGANISM: Homo sapiens
278 <400> SEQUENCE: 24
279 cgatcgactc tcagcaagcc tcaacgatcg      30
282 <210> SEQ ID NO: 25
283 <211> LENGTH: 27
284 <212> TYPE: DNA
285 <213> ORGANISM: Homo sapiens
287 <400> SEQUENCE: 25

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288 cgatcgtctc agcgggcctc acgatcg

27

RAW SEQUENCE LISTING ERROR SUMMARY
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Input Set : A:\01-18-07 seq list_ST25.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:9; N Pos. 13 ✓
Seq#:10; N Pos. 13 ✓
Seq#:12; N Pos. 13 ✓
Seq#:20; N Pos. 19 ✓
Seq#:21; N Pos. 19 ✓
Seq#:23; N Pos. 19 ✓

VERIFICATION SUMMARY

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Input Set : A:\01-18-07 seq list_ST25.txt

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L:105 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:246 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:270 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0